

Alignment View Print

	1 10 20
1Translation of 223122[residues 1-468]	R P L C S L S P R S C C W S O L S L S
DR5mature	- - - - - E S A L I T O Q D L A P
Consensus	x x x x x x x x S x x x x Q x x L x x
 30 40
1Translation of 223122[residues 1-468]	P N K T L P S R E W P H N K R G P A P Q
DR5mature	Q Q R A A P Q Q K R S S P S E G L C P
Consensus	x x x x x P x x x x x x x x G x x P x
 50 60
1Translation of 223122[residues 1-468]	R D C V H L D T I S Q K T V E I A S P A
DR5mature	G H H I S E D G R - D - - - - C I S C
Consensus	x x x x x D x x x x x x x x x x x x
 70 80
1Translation of 223122[residues 1-468]	N M D R T I A L T W N D L L F C L R C T
DR5mature	K Y G Q D Y S T H W N D L L F C L R C T
Consensus	x x x x x x x x x W N D L L F C L R C T
 90 100
1Translation of 223122[residues 1-468]	R C D S G E V E L S P C T T D Q K H S V
DR5mature	R C D S G E V E L S P C T T T R N T V C
Consensus	R C D S G E V E L S P C T T x x x x x x
 110 120
1Translation of 223122[residues 1-468]	S V R R R H L P G K K I L L R C A G S A
DR5mature	Q C E E G T F R - E E D S P E M C R K C
Consensus	x x x x x x x x x x x x x x x x x x

Alignment View Print

1Translation of 223122[residues 1-468] A Q G V P X D W M K V G E L Y T L E V T 130 140
 DR5mature R T G C P R G M V K V G D C T P W S D I

Consensus x x G x P R x x V K V G x x x x x x x x

1Translation of 223122[residues 1-468] S N V S T K N Q A S S E S 150 160
 DR5mature E C V H K E S G I I I G V T V A A V V L

Consensus x x V x x x x x x x x x x x x x x x x x x

1Translation of 223122[residues 1-468] 170 180
 DR5mature I V A V F V C K S L L W K K V L P Y L K

Consensus x x x x x x x x x x x x x x x x x x

1Translation of 223122[residues 1-468] 190 200
 DR5mature G I C S G G G G D P E R V D R S S Q R P

Consensus x x x x x x x x x x x x x x x x x x

1Translation of 223122[residues 1-468] 210 220
 DR5mature G A E D N V L N E I V S I L Q P T Q V P

Consensus x x x x x x x x x x x x x x x x x x

1Translation of 223122[residues 1-468] 230 240
 DR5mature E Q E M E V Q E P A E P T G V N M L S P

Consensus x x x x x x x x x x x x x x x x x x

Alignment View Print

1Translation of 223122[residues 1-468] 250 260

DR5mature G E S E H L L E P A E A E R S Q R R R L

Consensus x

1Translation of 223122[residues 1-468] 270 280

DR5mature L V P A N E G D P T E T L R Q C F D D F

Consensus x

1Translation of 223122[residues 1-468] 290 300

DR5mature A D L V P F D S W E P L M R K L G L M D

Consensus x

1Translation of 223122[residues 1-468] 310 320

DR5mature N E I K V A K A E A A G H R D T L Y T M

Consensus x

1Translation of 223122[residues 1-468] 330 340

DR5mature L I K W V N K T G R D A S V H T L L D A

Consensus x

1Translation of 223122[residues 1-468] 350 360

DR5mature L E T L G E R L A K Q K I E D H L L S S

Consensus x

Alignment View Print

..... 370 380
1Translation of 223122[residues 1-468]
DR5mature G K F M Y L E G N A D S A M S

Consensus x x x x x x x x x x x x x x x

Alignment View Print

	1 10 20	
2Translation of 223122[residues 2-469]	- - - - -	- - - - -
DR5mature	E S A L I T Q Q D L A P Q Q R A A P Q Q	
Consensus	x x x x x x x x x x x x x x x x x x	
 30 40	
2Translation of 223122[residues 2-469]	- - - - -	- - - - -
DR5mature	K R S S P S E G L C P P G H H I S E D G	
Consensus	x x x x x x x x x x x x x x x x x x	
 50 60	
2Translation of 223122[residues 2-469]	- - - - -	- - - - -
DR5mature	R D C I S C K Y G Q D Y S T H W N D L L	
Consensus	x x x x x x x x x x x x x x x x x x	
 70 80	
2Translation of 223122[residues 2-469]	- - - - -	- - - - -
DR5mature	F C L R C T R C D S G E V E L S P C T T	
Consensus	x x x x x x x x x x x x x x x x x x	
 90 100	
2Translation of 223122[residues 2-469]	- - - - -	- - - - - D
DR5mature	T R N T V C Q C E E G T F R E E D S P E	
Consensus	x x x x x x x x x x x x x x x x x x	
 110 120	
2Translation of 223122[residues 2-469]	P C A R C R R G P A V G L S V C S D H P	
DR5mature	M C R K C R T G C P R G M V K V G D C T	
Consensus	x C x x C R x G x x x G x x x x x D x x	



Alignment View Print

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. . . . . 130 . . . . . 140
2Translation of 223122[residues 2-469] T R P S S P A E S - - - - - G P T T
DR5mature P W S D I E C V H K E S G I I I G V T V

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Consensus x x x x x x x x x x x x x x x G x T x

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. . . . . 150 . . . . . 160
2Translation of 223122[residues 2-469] K E V Q P L R G I V S T W T P Y L R R R
DR5mature A A V V L I V A V F V C K S L L W K K V

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Consensus x x V x x x x x x x x x x x x x x x x x x

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. . . . . 170 . . . . . 180
2Translation of 223122[residues 2-469] R L H L L Q I W T G - - - - L H S L G M
DR5mature L P Y I K G I C S G G G G D P E R V D R

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Consensus x x x L x x I x x G x x x x x x x x x x

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. . . . . 190 . . . . . 200
2Translation of 223122[residues 2-469] T S F S A C A A P G V I Q V K W S V P A
DR5mature S S Q R P G A E D N V L N E I V S I L Q

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Consensus x S x x x x A x x x V x x x x x S x x x

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. . . . . 210 . . . . . 220
2Translation of 223122[residues 2-469] P P T R - - N T V C Q C E E G T F R E R
DR5mature P T Q V P E Q E M E V Q E P A E P T G V

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Consensus P x x x x x x x x x x x E x x x x x x x

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. . . . . 230 . . . . . 240
2Translation of 223122[residues 2-469] R F S D V P E V P H R V S Q X I G S R S
DR5mature N M L S P G E S E H L L E P A E A E R S

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Alignment View Print

Consensus	x x x x x x E x x H	x x x x A x x x R S
2Translation of 223122[residues 2-469]	V N - - - C T P W K	H R M C P Q R I R H
DR5mature	Q R R R L L V P A N	E G D P T E T L R Q
Consensus	x x x x x x x P x x	x x x x x z x x R x
2Translation of 223122[residues 2-469]	H H R S H	
DR5mature	C F D D F A D L V P	F D S W E P L M R K
Consensus	x x x x x x x x x x	x x x x x x x x x x
2Translation of 223122[residues 2-469]	L G L M D N E I K V	A K A E A A G H R D
DR5mature		
Consensus	x x x x x x x x x x	x x x x x x x x x x
2Translation of 223122[residues 2-469]	T L Y T M L I K W V	N K T G R D A S V H
DR5mature		
Consensus	x x x x x x x x x x	x x x x x x x x x x
2Translation of 223122[residues 2-469]	T L L D A L E T L G	E R L A K Q K I E D
DR5mature		
Consensus	x x x x x x x x x x	x x x x x x x x x x
2Translation of 223122[residues 2-469]	H L L S S G K F M Y	L E G N A D S A M S
DR5mature		

Alignment View Print

Consensus

x x

2Translation of 223122[residues 2-469] 370 380

DR5mature

Consensus



Alignment View Print

1 10 20
 3Translation of 223122[residues 3-467] T L V L V V A A V L L L V S A E S A L I
 DR5mature - - - - - E S A L I

Consensus x x x x x x x x x x x x x x E S A L I

. 30 40
 3Translation of 223122[residues 3-467] T Q Q D L A P Q Q R V A P Q Q K R S S P
 DR5mature T Q Q D L A P Q Q R A A P Q Q K R S S P

Consensus T Q Q D L A P Q Q R x A P Q Q K R S S P

. 50 60
 3Translation of 223122[residues 3-467] S E G L C P P G H H I S E D G R D C I S
 DR5mature S E G L C P P G H H I S E D G R D C I S

Consensus S E G L C P P G H H I S E D G R D C I S

. 70 80
 3Translation of 223122[residues 3-467] C K Y G Q D Y S T H L E P P F L L A L H
 DR5mature C K Y G Q D Y S T H W N D L L F C L R C

Consensus C K Y G Q D Y S T H x x x x x x x x x x

. 90 100
 3Translation of 223122[residues 3-467] Q V F R S G - - A K S L H H R P E T Q C
 DR5mature T R C D S G E V E L S P C T T T R N T V

Consensus x x x x S G x x x x S x x x x x x x x x

. 110 120
 3Translation of 223122[residues 3-467] V S A K K A P S G K E D S P E M C R K C
 DR5mature C Q C E E G T F R E E D S P E M C R K C

Consensus x x x x x x x x x x E D S P E M C R K C

Alignment View Print

3Translation of 223122[residues 3-467] 130 140
 DR5mature R T G C P R X L G Q G R I V H P G S D I
 R T G C P R G M V K V G D C T P W S D I

Consensus R T G C P R G x x x x x x x P x S D I

3Translation of 223122[residues 3-467] 150 160
 DR5mature E C V H K E S G I I I G V T V A A V V L
 E C V H K E S G I I I G V

Consensus E C V H K E S G I I I G V x x x x x x x

3Translation of 223122[residues 3-467] 170 180
 DR5mature I V A V F V C K S L L W K K V L P Y L K

Consensus x x x x x x x x x x x x x x x x x x

3Translation of 223122[residues 3-467] 190 200
 DR5mature G I C S G G G G D P E R V D R S S Q R P

Consensus x x x x x x x x x x x x x x x x x x

3Translation of 223122[residues 3-467] 210 220
 DR5mature G A E D N V L N E I V S I L Q P T Q V P

Consensus x x x x x x x x x x x x x x x x x x

3Translation of 223122[residues 3-467] 230 240
 DR5mature E Q E M E V Q E P A E P T G V N M L S P

Consensus x x x x x x x x x x x x x x x x x x

Alignment View Print

3Translation of 223122[residues 3-467] 250 260

DR5mature G E S E H L L E P A E A E R S Q R R R L

Consensus x

3Translation of 223122[residues 3-467] 270 280

DR5mature L V P A N E G D P T E T L R Q C F D D F

Consensus x

3Translation of 223122[residues 3-467] 290 300

DR5mature A D L V P F D S W E P L M R K L G L M D

Consensus x

3Translation of 223122[residues 3-467] 310 320

DR5mature N E I K V A K A E A A G H R D T L Y T M

Consensus x

3Translation of 223122[residues 3-467] 330 340

DR5mature L I K W V N K T G R D A S V H T L L D A

Consensus x

3Translation of 223122[residues 3-467] 350 360

DR5mature L E T L G E R L A K Q K I E D H L L S S

Consensus x

Alignment View Print

• 3Translation of 223122[residues 3-467] 370 380
DR5mature G K F M Y L E G N A D S A M S
•
Consensus x x x x x x x x x x x x x x

